



SEQUENCE LISTING

- <110> Quint, Wilhelmus Van Doorn, Leendert
- <120> PROBES, METHODS AND KITS FOR DETECTION
 AND TYPING OF HELICOBACTER PYLORI NUCLEIC ACIDS IN
 BIOLOGICAL SAMPLES
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- <140> 10/035,978
- <141> 2001-12-21
- <150> 09/284,725
- <151> 1999-04-16
- <150> EP 97870133.2
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- <150> EP 96870131.8
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<210> 151
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<223> Helicobacter pylori vacA nucleic acid sequence
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<210> 153
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<221> misc_feature
<222> 27, 34, 53, 55, 76, 82
<223> n = A, T, C \text{ or } G
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<223> Helicobacter pylori vacA nucleic acid sequence
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<211> 105
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<223> Helicobacter pylori vacA nucleic acid sequence
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<211> 105
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<210> 169
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<222> 82
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<221> misc_feature
<222> 82
<223> n = A, T, C \text{ or } G
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<210> 170
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<211> 105
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<210> 174
<211> 105
<212> DNA
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<400> 174
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<210> 175
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<222> 26, 27
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<211> 105
<212> DNA
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<221> misc_feature
<222> 82, 101, 102, 103, 104
<223> n = A, T, C \text{ or } G
<221> misc feature
<222> 82, 101, 102, 103, 104
<223> n = A,T,C or G
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<211> 105
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<211> 105
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<211> 105
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<211> 105
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<221> misc_feature
<222> 7, 27, 34, 55, 82
<223> n = A,T,C or G
<221> misc_feature
<222> 7, 27, 34, 55, 82
<223> n = A, T, C or G
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<223> Helicobacter pylori vacA nucleic acid sequence
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<210> 194
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ctaaacacta gcactttgga ttttagcggc gttacagaca aagtcaatat caacaagctc 180
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accogtgttc agagttttgg gcaatacact atttttggcg aaaatatagg cgataagtct 300
cgcattggtg tcgttagttt gcaaactggc tatagcccgg cctattctgg gggcgttact 360
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ctaaacacta gtgctttgga ttttagcggc gttacagaca aagtcaatat caacaagctc 180
actacatctg ccactaatgt gaacattaaa aactttgaca ttaaggaatt agtggttaca 240
accogagtto aaagttttgg gcaatacact atttttggcg aaaatatagg cgataagtct 300
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ctaaacacta gtgctttgga ttttagcggc gttacagaca aagtcaatat caacaagctc 180
actacatetg ccactaatgt gaacattaaa aactttgaca ttaaggaatt agtggttaca 240
accegagete aaagetetegg geaatacact atteteggeg aaaatatagg egataageet 300
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ctaaacacta gtgctttgga ttttagcggc gttacagaca aagtcaatat caacaagctc 180
actacatctg ccactaatgt gaacattaaa aactttgaca ttaaggaatt agtggttaca 240
accogagtto aaagttttgg goaatacact atttttggcg aaaatatagg cgataagtot 300
cgcattggtg tcgtgagttt gcaaacggga tatagcccgg cctattctgg gggcgttact 360
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actacatctg ccactaatgt gaacattaaa aactttgaca ttaaggaatt agtggttaca 240
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